

are inserts absent in htau23 and htau24 so that these molecules have only 6 ser-pro motifs. The following radioactive tryptic peptides were found:

E1
cont.

29-49:	KDQGGYTMHQDQEGDTDAGKLKES _p PLQ (SEQ ID NO: 31)
191-209:	SGDRSGYSS _p P _p GSPGTPGSR (SEQ ID NO: 32)
231-240:	TPPKS _p PSSAK (SEQ ID NO: 33)
396-406:	SPVVSGDTS _p PR (SEQ ID NO: 34)
386-406:	TDHGAEIVYKS _p PVVSGDTS _p PR (SEQ ID NO: 35)
407-428:	HLSNVAATGA IDMVDS _p PQLATL (SEQ ID NO: 36)
260-266:	IGS _p TENL (SEQ ID NO: 37) --

Please replace the first full paragraph at page 60, with the following rewritten paragraph:

E2

-- Since it was suspected that the two KSP motifs were phosphorylated by the kinase, it was tried to prove this directly. Radioactively labeled tryptic peptides of htau34 were identified by HPLC and protein sequencing, and phosphorylated residues were determined. There are two major phosphorylated tryptic peptides in these regions; peptide 1 (T231-K240, TPPKS_pPSSAK (SEQ ID NO.: 33)) contains the first KSP motif, phosphorylated at S235, peptide 2 (T386-R406, TDHGAEIVYKS_pPVVSGDTS_pPR (SEQ ID NO: 35)) contains the second KSP site, phosphorylated at S396 and S404. S416, the single phosphorylation site of CaM kinase described earlier (Steiner et al., EMBO J. 9 (1990), 3539-3544, S405 in the numbering of htau23 used earlier) is not phosphorylated by the kinase used here. --

IN THE SEQUENCE LISTING

Please replace the sequence listing as originally filed with the substitute sequence listing (pages 1-10) submitted herewith.

II. REMARKS

Accompanying the paper copy of the substitute Sequence Listing is a computer-readable version of the sequence listing as well as a "Statement Pursuant to 37 C.F.R. §1.825(a) and (b)." Applicant submits that no new matter has been added via the submission of the Sequence Listing or amendment.